



Cite this: *CrystEngComm*, 2021, 23, 2027

Received 9th January 2021,
Accepted 26th February 2021



DOI: 10.1039/d1ce00041a

rsc.li/crystengcomm

We report on nanoindentation data for two pairs of polymorphic compounds of *p*-aminobenzoic acid (*p*ABA) and *p*-nitrobenzoic acid (*p*NBA). Variations of 60% or higher are observed in both stiffness and hardness data between the two forms of each pair. We interpret these observations through variations of periodic bond chains (PBCs), *i.e.* variations in continuous intermolecular interactions strength and nature. Aided with hardness data of polymorphs from the literature, we then further assess how PBCs can be used to assess and compare mechanical properties of polymorphs. We define a new parameter, *s*-PBC, which quantifies the strongest PBC interaction that needs breaking so as to allow a switch of the PBC network from 3D to 2D/1D. We found that higher values of *s*-PBCs, normalised by the mean molecular volumes, usually led to higher values of mechanical properties. Although not infallible, this trend seemed to hold well when comparing polymorphs of a given compound, but not so much when comparing across different compounds.

Mechanical properties impact the manufacturing routes of molecular materials since they strongly determine the processing steps required.^{1–3} The mechanical properties of solids are intimately related to their constituent molecules and chemistry as well as their short and long-range arrangements in the solid state. Notably, the ability of many molecules to crystallise into different crystal structures, known as polymorphs,⁴ is well known as well as the consequences of polymorphism in a material's physicochemical properties^{5–7} and intellectual property (IP).⁸ Due to the strong economic impact of polymorphism on IP, as well as the fact that variations of physical properties across polymorphic forms can hinder or enable manufacturing

Impact of polymorphism on mechanical properties of molecular crystals: a study of *p*-amino and *p*-nitro benzoic acid with nanoindentation†

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(potentially leading to substantial money and time loss or gained), increasing efforts have been directed in the past decades towards developing computational crystal structure and polymorph prediction methods.^{9,10} Beyond the prediction of structure, much work has been focused in recent years on achieving a qualitative prediction of the mechanical behaviour of crystalline molecular crystals such as shearing, bending or brittleness¹¹ – sometimes between polymorphs.¹² Further work is needed, however, on quantitative predictions of mechanical properties. To achieve this, we still need to accurately measure stiffness and hardness of a significant number of molecular crystals and their polymorphs so as to provide a significant dataset against which theoretical models can be compared. Recent developments in the area of nanoindentation^{13,14} offer opportunities to remedy this.

Following a first study of nanoindentation of a series of *p*-derivatives of benzoic acid crystals,¹⁵ we investigate here the impact of polymorphism on mechanical properties of *p*-nitrobenzoic acid (*p*NBA) and *p*-aminobenzoic acid (*p*ABA). Both compounds crystallise into various polymorphs, including forms I (CSD refcode: NBZOAC03 (ref. 16)) and II (NBZOAC14 (ref. 17)) for *p*NBA and forms α (AMBAC06 (ref. 18)) and β (AMBAC04 (ref. 19)) for *p*ABA. Both pairs of polymorphs are related enantiotropically with transition temperatures of 51 °C (ref. 20) for *p*NBA (*p*NBA-I being the low temperature form) and 13.8 °C for *p*ABA (β -*p*ABA being the low temperature form).^{21,22} *p*NBA crystallises readily into its form I at room temperature and into its form II at temperatures above 51 °C.²⁰ Both polymorphs of *p*ABA and *p*NBA, once crystallised and removed from their crystallisation solution, are also reported to not convert to another form at room temperature over time.^{20,23} *p*ABA crystallises into its α -form from all temperatures and most solvents²⁴ except for water at low supersaturation levels from which β -*p*ABA is crystallised. Using mechanical data derived from nanoindentation experiments on these two pairs of polymorphs, we rationalise the observed variations in stiffness and hardness between polymorphs from observed

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† Electronic supplementary information (ESI) available. See DOI: 10.1039/d1ce00041a



differences in intermolecular interactions and crystal packing, and we compare the observed variations with those reported for other polymorphic systems.

Comparing mechanical data across different polymorphs can be challenging because mechanical properties of single crystals are highly anisotropic thus they depend on the crystal orientation (the hkl values of the faces indented).²⁵ In our previous study of mechanical properties of benzoic acid derivatives measured by nanoindentation, we were able to compare stiffness and hardness data across different compounds because they all crystallised with common crystal packing features (consisting of hydrogen bonded dimers and infinite aromatic stacks) and they were indented along equivalent crystal directions.¹⁵ Since crystal structure and packing change in different polymorphs, it is not obvious which crystal orientations, if any, could enable the most relevant comparison of mechanical properties between different forms. In *p*ABA, for example, the α -form consists of continuous aromatic stacking interactions along the b -axis and discrete, 0D $R_2^2(8)$ hydrogen bonded carboxylic acid dimers. Instead, the β -form is dominated by continuous hydrogen bonds between carboxylic and amine groups and 0D inverted aromatic stacked dimers. In *p*NBA, forms I and II have some similarities, including discrete $R_2^2(8)$ carboxylic acid dimers and continuous aromatic stacking. Furthermore, the dominant crystal face for both *p*NBA forms I and II renders the aromatic stacking interactions orthogonal to the indentation direction (*i.e.* parallel to the indenter stage). *p*NBA forms I and II in fact share a common layer of molecules whereby the carboxylic acid dimers and aromatic stacks are arranged identically (presented in ESI†). These

layers, however, pack in different ways by means of different symmetry operations (form I crystallises in $A2/a$ whilst form I $P2_1/c$). Micrographs of single crystals of *p*ABA and *p*NBA polymorphs with their respective face indexing are shown in Fig. 1 (data for α -*p*ABA and *p*NBA-I were retrieved from a previous study).¹⁵ We note that two faces are highlighted for α -*p*ABA in Fig. 1, (200) and (002), because they are crystallographically and structurally identical and thus they cannot be differentiated. Thus, on indenting different crystals of α -*p*ABA, some would have been indented on the (200) face and others on the (002) but no differences are observed in the mechanical behaviour because of the structural identity between these faces.

Typical load-depth (P - d) curves measured with nanoindentation for the α -*p*ABA and β -*p*ABA are presented in Fig. 2a and for *p*NBA-I and *p*NBA-II in Fig. 2b. For *p*ABA, the experiments were done in load-controlled mode. At maximum load (for both forms, $P_{MAX} = 5000 \mu\text{N}$), the final depth of penetration reached for β -*p*ABA was just 400 nm whilst it was 600 nm for α -*p*ABA. Many sudden bursts (pop-ins) occurred in the loading section of the P - d curve on β -*p*ABA (Fig. 2a, red curve, black arrows) whilst those were hardly distinguishable in α -*p*ABA (only very light discontinuities were observed on loading). For *p*NBA, the experiments were done in displacement-controlled mode. At maximum depth of penetration, h_{MAX} , the maximum load required for form I was twice higher than that of form II. As in α -*p*ABA, the loading section of form I *p*NBA has very light discontinuities but cannot be designated as real “bursts”. In form II *p*NBA, however, both very small pop-ins and some discontinuities are observed. Of particular interest, the unloading sections of all curves in Fig. 2

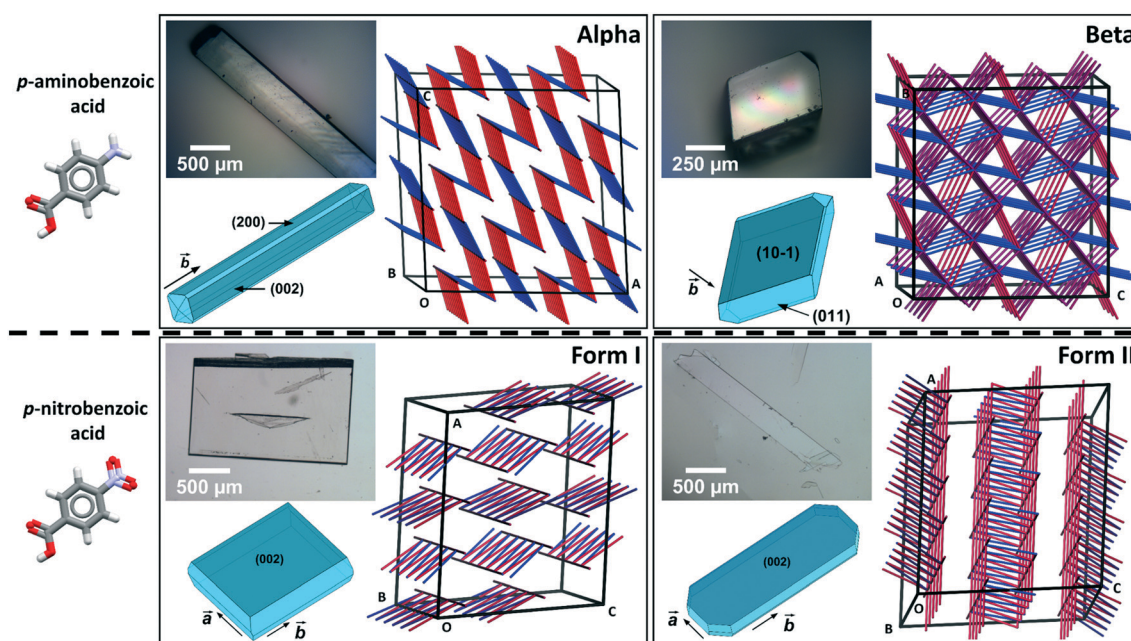


Fig. 1 (Upper) Molecule of *p*ABA, micrographs and face indexing of its alpha and beta polymorph crystals with their respective PBCs. (Lower) Molecule of *p*NBA, micrographs and face indexing of its forms I and II crystals with their respective PBCs. Indented faces are shown in darker blue in the face indexed morphology graph.





Fig. 2 (a) Typical load-depth curves obtained from indenting α -*p*AABA (black curve) and β -*p*AABA (red curve) and (b) typical load-depth curves obtained from indenting *p*NBA form I (black curve) and form II (red curve). Indents on *p*AABA were carried out in load-controlled mode up to $P_{\text{MAX}} = 5$ mN, hence polymorphs can be readily compared using the final depth reached. For *p*NBA, indents were carried out in displacement-control mode up to $h_{\text{MAX}} = 425$ nm, thus the maximum load reached on each polymorph should be compared.

are “pop-out” free. “Pop-out” events are sudden burst towards lower penetration depth upon indentation pressure release and are characteristic of pressure-induced polymorphic transformation.^{26,27} Here, their absence are suggesting that no polymorphic transformation occurred upon pressure released, which may suggest that no polymorphic transformation occurred at all upon indentation and that E and H values extracted correspond to the true values of the polymorphs.

Table 1 shows the derived elastic modulus (using a Poisson ratio of 0.3 for *p*AABA and *p*NBA, reported as a typical value for molecular single crystals²⁵) and hardness data obtained from indenting the polymorphs of these two systems together with other data on polymorphs retrieved from the literature. In *p*AABA, the elastic modulus E and the hardness H are *ca.* 60% and 130% higher respectively for β -*p*AABA than for α -*p*AABA. In *p*NBA, E and H are *ca.* 60% and 80% higher respectively for form I than for form II. The periodic bond chains (PBCs) networks for these crystal structures are shown Fig. 1. PBCs illustrate the network structure of the intermolecular interactions in the crystals. Using a PBC cut-off value of -13.1 kJ mol⁻¹, *p*NBA-I has a 1D PBC network, *p*NBA-II and α -*p*AABA have 2D PBC networks whilst β -*p*AABA has a 3D PBC network. The crystal structure of β -*p*AABA is dominated by an infinite head to tail (between the carboxylic acid and the amino group) hydrogen bonds (HBs) which further assemble into a 3D-HB PBC network. In β -*p*AABA, each molecule is involved in 4 HBs. The 3D PBCs network based on strong HBs (-19 kJ mol⁻¹ is the weakest interaction in the β -*p*AABA 3D network) leads to higher mechanical properties for β -*p*AABA compared to the other systems. We notice that the loading section of the load-depth trace for β -*p*AABA has marked pop-ins in the load-depth curve. These pop-ins could be associated with possible fracturing,

although none was observed at the surface of β -*p*AABA single crystals upon indentation, as shown on AFM images in Fig. 3. Images of indents on the four systems revealed crack-free indents (see ESI†). In summary for *p*AABA, β -*p*AABA with 3D PBCs is stiffer and harder than α -*p*AABA with 2D PBCs. For *p*NBA, form I with 1D PBCs is stiffer and harder than form II, however with 2D PBCs.

Table 1 collects E and H data for our polymorphs of *p*AABA and *p*NBA as well as for sulfathiazole²⁸ (STZ), curcumin²⁸ (CCM), a fluorinated amide²⁹ (FIA), famotidine³⁰ (FAM), febuxostat³¹ (FEB) and felodipine³² (FEL) polymorphs reported by other groups (all measured by nanoindentation with a Berkovich tip). Table 1 also summarises on the PBC dimensionality of all those crystal structures (using a cut-off value of -13.1 kJ mol⁻¹) as well as the PBC value switching the networks from 3D to 2D/1D (s-PBC), the nature of the s-PBC interaction, the (hkl) of the strongest 2D plane with its rugosity, and the direction of easiest shear for all those systems. We note that the s-PBC is the PBC interaction that needs breaking so as to allow a switch of the PBC network from 3D to 2D or 1D network. The value is given as positive since the interaction needs breaking. We explore this value here since once this s-PBC is broken this would allow for some slip systems. Thus, less stabilising s-PBCs will be easier to break and give rise to slip planes, thus this may be related to softer material responses. Additionally, the strongest 2D layer remaining after breaking the s-PBC is identified with its (hkl) value and its rugosity calculated using the Bryant, Maloney and Sykes method.³³

Two main observations can be drawn from the data in Table 1. First, there is a tendency for hard materials to have 3D PBCs and higher absolute values of s-PBCs. A molecular level identification of these s-PBC (Table 1) shows that PBCs



Table 1 *E* and *H* data (averages, standard deviation is given for data from this contribution) measured by nanoindentation on *p*NBA forms I and II and *p*A_{BA} forms α and β (this work). Data reported in the literature of mechanical properties of single component polymorphs are also included. Space group (SG) and *hkl* indices of the indented face are also given, along with the ratio highest/lowest for both data of each polymorphic system. Information on *s*-PBCs is also given. For the calculation of the PBC dimension (PBC dim), an energy cut-off of -13.1 kJ mol $^{-1}$ was used

System ^a	SG	Indent. face	CSD refcode	PBC dim.	<i>s</i> -PBC (kJ mol $^{-1}$)	Nature of <i>s</i> -PBC	(<i>hkl</i>) 2D layer, rugosity (Å)	Easiest shear direction ^b	<i>E</i> (\pm stdv) (GPa)	<i>H</i> (\pm stdv) (MPa)	<i>E</i> ratio ^c	<i>H</i> ratio ^c
<i>p</i> A _{BA} - α	<i>P</i> ₂ ₁ / <i>n</i>	(200)	AMBNA06	2D	13.8	NH \cdots O=C	1D, -0.4	<i>b</i> -axis [t]	7.17 \pm 0.25	221 \pm 17	1.6	2.3
<i>p</i> A _{BA} - β ^d	<i>P</i> ₂ ₁ / <i>n</i>	(10 $\bar{1}$)	AMBNA04	3D	19.8	NH \cdots O=C	(010), -1.0	<i>a</i> -axis [t]	11.38 \pm 0.32	512 \pm 25		
<i>p</i> NBA-I ^d	<i>A</i> 2/ <i>a</i>	(002)	NBZOAC03	1D	9.8	Other	(001), 0.0	<i>b</i> -axis [t]	9.19 \pm 0.48	322 \pm 22	1.6	1.8
<i>p</i> NBA-II	<i>P</i> ₂ ₁ / <i>c</i>	(002)	NBZOAC14	2D	9.8	Other	(001), 0.4	<i>a</i> -axis [t]	5.88 \pm 0.25	182 \pm 10		
STZ-I ²⁸	<i>P</i> ₂ ₁ / <i>n</i>	(100)	SUTHAZ28	3D	22.8	NH \cdots O=S	(001), -2.9	<i>a</i> -axis [i]	10.01	356	2.0	3.0
STZ-II ²⁸	<i>P</i> ₂ ₁ / <i>c</i>	(100)	SUTHAZ24	3D	31.0	NH \cdots O=S	(100), 0.2	<i>b</i> -axis [t]	20.44	1080		
STZ-III ²⁸	<i>P</i> ₂ ₁ / <i>c</i>	(100)	SUTHAZ25	3D	30.4	NH \cdots N=	(100), 0.2	<i>b</i> -axis [z']	16.42	704		
STZ-IV ²⁸	<i>P</i> ₂ ₁ / <i>n</i>	(10 $\bar{1}$)	SUTHAZ26	3D	17.5	NH \cdots N=	(10 $\bar{1}$), 0.1	<i>b</i> -axis [s]	17.31	881		
CCM-I ²⁸	<i>P</i> ₂ / <i>n</i>	(001)	BINMEQ02	3D	26.1	Other	(101), -0.2	<i>b</i> -axis [i]	11.15	432	2.0	1.3
CCM-II ²⁸	<i>Pca</i> 2 ₁	(100)	BINMEQ08	3D	22.0	OH \cdots O-	(100), -3.8	<i>b</i> -axis [z't]	5.68	341		
CCM-III ²⁸	<i>Pbca</i>	(001)	BINMEQ07	3D	21.8	OH \cdots O-	(001), -3.5	<i>b</i> -axis [g]	5.60	330		
FLA-I ²⁹	<i>P</i> ₂ ₁	(001)	VAPMEH	2D	7.0	Aromatic	(001), 0.6	<i>b</i> -axis [t]	8.36	347	1.4	2.0
FLA-II ²⁹	<i>P</i> 1	(001)	VAPMEH01	2D	6.1	Aromatic	(001), 0.5	<i>a</i> -axis [t]	6.17	172		
FAM-A ³⁰	<i>P</i> ₂ ₁ / <i>c</i>	(100)	FOGVIG04	3D	27.1	NH \cdots N=	(001), -1.9	<i>b</i> -axis [t]	22.6	1580	1.2	1.9
		(001)							20.1	1330		
FAM-B ³⁰	<i>P</i> ₂ ₁ / <i>n</i>	($\bar{1}$ 01)	FOGVIG05	3D	16.9	Other	($\bar{1}$ 01), -0.2	<i>b</i> -axis [t]	19.5	840		
FEB-Q ³¹	<i>P</i> ₂ ₁ / <i>c</i>	NR	HIQQAB	3D	24.9	Other	(010), -2.2	<i>a</i> -axis [t]	2.5	214	1.8	1.4
FEB-H1 (ref. 31)	<i>P</i> 1	NR	HIQQAB02	3D	20.8	Other	(010), -0.6	<i>a</i> -axis [z']	4.5	310		
FEL-I ³²	<i>P</i> ₂ ₁ / <i>c</i>	(100)	DONTIJ	3D	32.7	NH \cdots O	1D, -0.5	<i>a</i> -axis [t]	12.8	553	2.1	2.2
FEL-II ³²	<i>C</i> 2/ <i>c</i>	(100)	DONTIJ04	3D	26.8	Other	(001), -0.4	<i>b</i> -axis [i]	6.2	260		
FEL-III ³²	<i>P</i> ₂ ₁ / <i>n</i>	(10 $\bar{1}$)	DONTIJ02	3D	26.0	Other	(101), -1.1	<i>b</i> -axis [t]	10.5	451		

^a STZ = sulfathiazole; CCM = curcumin; FLA = fluorinated amide; FAM = famotidine; FEB = febuxostat; FEL = felodipine. ^b This was estimated visually. Since many of the 2D layers are corrugated, shear along the direction of the ridges would be easier than against the ridges (see Bryant, Maloney and Sykes). The symmetry operation applied upon shear of one molecular position along this direction is also given. The symmetry relationships are: t = translation, i = inversion, g = glide plane, s = screw and z' = no symmetry due to $Z' > 1$. ^c Highest over lowest. ^d This work.

involving hydrogen bond interactions are usually stronger in nature than *s*-PBC involving other type of van der Waals dominated interactions or aromatic interactions. However, when molecules are larger (*i.e.* FEL), other type of interactions may become as important as hydrogen bonds. Usually, across a family of polymorphs, *s*-PBC tend to be of a similar nature though not always. Second, although we observed this trend for most of the compounds in the table, still some polymorphs may not obey this trend, such as FEB

or STZ-IV. These correlations are complex since there are many plausible mechanisms of plastic deformation which would affect the resulting hardness of the material (*i.e.* anisotropy, plasticity due to impurity or imperfection profile, or plastic deformation initiated from the smoothest plane instead of the weakest).

Whilst some trends in stiffness and hardness may be derived across isomechanical groups of crystals with similar packings and interactions (for example between *E* and melting points^{15,34} or lattice energies per unit of volume and hardness³⁵), such correlations may be harder to achieve for polymorphic structures since the crystal packings can be very different. In previous studies of nanoindentation on polymorphs (including those in Table 1), it has been shown that there is a large variation in mechanical properties between polymorphs. The ratio of *E* and *H* in polymorphs can lie somewhere between 1.2–2.0 for *E* and 1.3–3.0 for *H*. These large variations cannot match the small differences in melting points or lattice energies (see ESI†) since these are both very similar in polymorphs. First principle simulations, however, have been promisingly used for the direct computation of stiffness matrices for various polymorphic systems.^{36–38} Prediction of hardness is a much more complex exercise.

When comparing data of polymorphs within individual systems in Table 1, rather than comparing all data altogether,



Fig. 3 Height images (8 μ m scan size) of indents made on a single crystal of a) α -*p*A_{BA} and b) β -*p*A_{BA}, acquired using atomic force microscopy (AFM) in PeakForce QNM mode in air.



a clear trend appears. Fig. 4 shows the hardness of every system in Table 1 as a function of their s-PBC normalised by their mean molecular volume Ω . Values for polymorphs of different compounds are circled separately. For all systems but one (FEB) and mostly all polymorphs except for STZ-IV, there is a clear tendency for higher values of nanoindentation hardness to lead to higher values of s-PBC density. This is somewhat expected because higher values of s-PBC mean stronger 3D-networks with no easy slip-system and thus harder materials. The s-PBC model, however, is a very simple model based on PBCs also known as energy frameworks.³⁹ The novelty here is that the identification of the s-PBC allows for a quantification of the slip resistance. We note that for systems where correlations fail, the 2D layers may have different degrees of corrugation or the shear direction may require of different symmetry operations. The easiest shear direction was assumed as the direction of propagation of ridges for corrugated layers. Thus, a highly corrugated layer may be hard to shear against the corrugation but can be easily sheared along the propagation of ridges and valleys.³³ Here, the symmetry is important. Shear along a direction related by translation may result in the breaking and reforming of interactions as the molecules shear (slip). Shear along a direction related by other symmetry operations, however, may result in molecules occupying a lattice site with the wrong orientation which in turn may cause strains and repulsions and perhaps even cracks or brittle behaviour. We note that the systems that tend to fail the correlations (FEB and STZ-IV) do have differing layer rugosities and differing symmetry operations along the easiest shear directions.

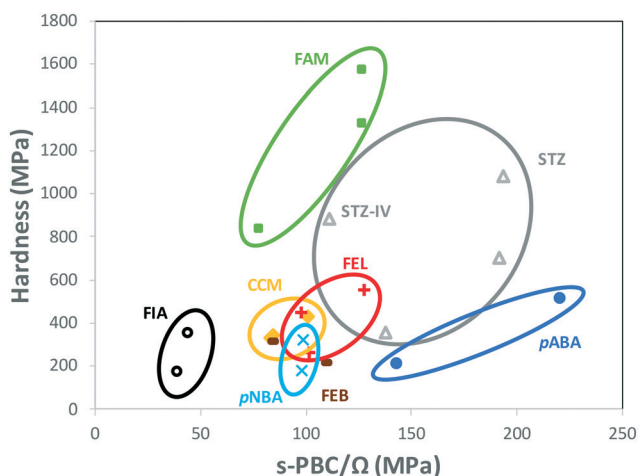


Fig. 4 Hardness measured with nanoindentation on various polymorphic systems as a function of the s-PBC normalised by the mean molecular volume Ω . Circles contain data for individual polymorphic systems. The general trends per individual system show an increase in hardness with s-PBC, but such trend is less obvious when considering all systems together. We note that febusostat (FEB, brown symbols) is the only system for which the trend shows that lower hardness is obtained at higher value of s-PBC/ Ω .

This finding is encouraging since the calculation of s-PBC is straight forward thus would provide a qualitative assessment of which polymorph would be expected to be harder. Deriving a quantitative tool, however, is much more challenging. The data spreads differently in Fig. 4 for different systems. For example, values of hardness for FIA and *p*ABA range from 172 to 512 MPa and are found respectively at the lowest and the highest values of s-PBC/ Ω . These observations should be taken carefully since there is still a lack of data available in the literature of single-component polymorphs to establish high quality, qualitative (and quantitative) predictions of mechanical properties of polymorphs.

In conclusion, we have reported here on mechanical properties measured by nanoindentation on two polymorphic compounds, *p*ABA and *p*NBA. We observed strong variations in *E* and *H* between polymorphic forms, in particular for *p*ABA which sees an increase of a factor of 2.3 between the hardness of its α and β forms. This particular variation can be quickly assessed by means of variations between s-PBCs of the forms. Using these new data together with previous literature data on hardness of polymorphs, we attempted to establish trends among mechanical properties. We found that hardness data can be, in some of these cases, qualitatively estimated for polymorphs of individual compounds using s-PBC/ Ω . This is encouraging since this simple method can be used for the qualitative prediction of relative hardness of polymorphs in, say, a computer-generated polymorphic landscape. However, there is a limited amount of data available on polymorphs of molecular compounds thus further experimental and computational studies are required to validate these trends. The roots of this model lie in the well-studied *energy frameworks*³⁹ which has been used by others to understand mechanical properties of different crystals forms³⁹ and also polymorphs:⁴⁰ s-PBC is just the interaction limit that turns the energy framework from 3D to 2D/1D. This is indeed a very simple model and many aspects of the system are not taken into consideration including molecular flexibility⁴¹ and its impact in mechanical behaviour, impact of temperature^{42,43} (calculations are done at 0 K) or interlocking of the plausible slip planes.³³ We see this model as a qualitative utility which should be of course taken with caution since the origin of the hardness of materials is indeed very complex.

Experimental section

Single crystals of *p*NBA (>98%, Fluka Analytical) forms I and II were produced by the slow evaporation of a saturated isopropanol (puriss. >99.5%, Merck) solution at RT and an acetonitrile (>99.9%, HPLC grade, Fisher Chemical) solution at 70 °C respectively. Crystals of α -*p*ABA were grown from slow cooling crystallisation experiments of supersaturated acetonitrile solutions (50 to 25 °C, -1 °C h^{-1} , supersaturation ratio $S = 1.6$). Large crystals of β -*p*ABA were obtained by seeding a slightly saturated aqueous solution with ~ 100 μm β -*p*ABA seeds. The solution was left to evaporate for 5–10 days for the seeds to grow and reach a size of *ca.* 0.5–1 mm.



Crystals were viewed under crossed polarised light (Axioplan 2 microscope, Zeiss, Jena, Germany) and only single crystals were retained whilst twinned crystals were discarded. Representative single crystals were mounted on a goniometer on a single crystal X-ray diffractometer (XCalibur-2, Rigaku Oxford Diffraction, Tokyo, Japan) with a MoK α X-ray source ($\lambda = 0.71073 \text{ \AA}$), and 360 successive images were taken of the crystals while rotating it. X-ray diffraction measurements were carried out, and using the software CrysAlisPro⁴⁴ (version 171.40.14d, Rigaku Oxford Diffraction), the unit cell parameters were verified to match those in the crystals' respective CSD refcode (see introduction) and face indexing was carried out, attributing their respective Miller indices to the different faces of the crystals studied.

Nanoindentation experiments were carried out on 7 to 10 single crystals per polymorph studied. Crystals were mounted only if their crystal habit clearly matched that assessed *via* face indexing and indented only if their surface tilt was lower than 3%. On each crystal, 9 indents were carried out using a Hysitron Ti950 Triboindenter (Bruker Nanomechanical Test Instruments, Minneapolis, USA) with a diamond Berkovich tip at a constant temperature of $23.1 \pm 0.5 \text{ }^\circ\text{C}$. Reduced elastic modulus and hardness data were extracted using the unloading section of the load-depth curves as described in ISO 14577.⁴⁵ Indents were first carried out at various loads or depths of penetration to determine the minimum load or depths at which minimal variation in the data was observed, as described in a previous study.¹⁴ Indents on *p*NBA crystals (both forms) were carried out in displacement-control mode, with a 5 second loading, a 2 second holding at $h_{\text{MAX}} = 425 \text{ nm}$, and a 5 second unloading LF sections. For *p*ABA, indents were carried out in load-control mode, with 5 seconds used for loading, holding and unloading, with $P_{\text{MAX}} = 5 \text{ mN}$. The distance between indents was $40 \text{ }\mu\text{m}$.

Periodic bond chains (PBCs) were computed from the crystal structures retrieved from the CSD using Material Studio 2019 (ref. 46) (Dassault Systèmes, BIOVIA) with the COMPASSII forcefield (and its atomic charges). The cut-off value of the PBC was increased progressively until the structure went from a 3D to a 2D/1D structure. This reveals the *s*-PBC value. Crystal structures were geometry optimised by relaxing all atomic positions in the unit cell while allowing for unit cell parameters to optimise prior to PBC calculations.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

The authors thank the University of Manchester and Hoffmann-La Roche for financial support. This work was supported by the Henry Royce Institute for Advanced

Materials, funded through EPSRC grants EP/S009493/1, EP/R00661X/1, EP/P025021/1, and EP/P025498/1.

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